BIOTECHNOLOGY
SYSTEMS
BRANCH

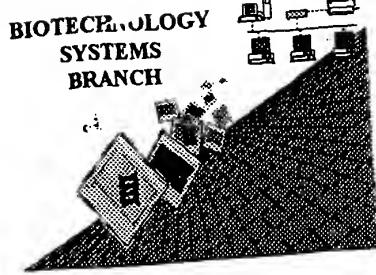# RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _09/659,983B_

Source: _OIPE_

Date Processed by STIC: _6/5/2001_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: **patin21help@uspto.gov** or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: **patin3help@uspto.gov** or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
http://www.uspto.gov/web/offices/pac/checker

RAW SEQUENCE LISTING   DATE: 06/05/2001
PATENT APPLICATION: US/09/659,983B   TIME: 14:22:06   Does Not Comply
Corrected Diskette Needed

Input Set : **A:\PTO.txt**
Output Set: **C:\CRF3\06052001\I659983B.raw**

pp 2,6

```
  3 <110> APPLICANT: Meloen, Robert Hans
  4        Oonk, Hendrica Berendina
  6 <120> TITLE OF INVENTION: An Improved Peptide, Immunogenic Composition and Vaccine or
Medical
  7        Preparation, a Method to Immunise Animals Against the Hormone LHRH, and Analogs
  8        of the LHRH Tandem Repeat Peptide and their Use as Vaccine
 10 <130> FILE REFERENCE: 2183-4518US
 12 <140> CURRENT APPLICATION NUMBER: 09/659,983B
 13 <141> CURRENT FILING DATE: 2001-05-12
 15 <150> PRIOR APPLICATION NUMBER: US 09/274,048
 16 <151> PRIOR FILING DATE: 1999-03-22
 18 <150> PRIOR APPLICATION NUMBER: US 08/981,557
 19 <151> PRIOR FILING DATE: 1995-06-07
 21 <150> PRIOR APPLICATION NUMBER: PCT/NL96/00223
 22 <151> PRIOR FILING DATE: 1996-06-06
 24 <150> PRIOR APPLICATION NUMBER: US 08/447,298
 25 <151> PRIOR FILING DATE: 1995-06-07
 27 <150> PRIOR APPLICATION NUMBER: US 08/476,013
 28 <151> PRIOR FILING DATE: 1995-06-07
 30 <160> NUMBER OF SEQ ID NOS: 13
 32 <170> SOFTWARE: PatentIn version 3.0
 34 <210> SEQ ID NO: 1
 35 <211> LENGTH: 10
 36 <212> TYPE: PRT
 37 <213> ORGANISM: Sus scrofa
 39 <220> FEATURE:
 40 <221> NAME/KEY: PEPTIDE
 41 <222> LOCATION: (1)..(1)
 42 <223> OTHER INFORMATION: X=pyroglutamic acid
 45 <220> FEATURE:
 46 <221> NAME/KEY: PEPTIDE
 47 <222> LOCATION: (10)..(10)
 48 <223> OTHER INFORMATION: X=Gly-NH2
 51 <400> SEQUENCE: 1
 53 Xaa His Trp Ser Tyr Gly Leu Arg Pro Xaa
 54 1               5                   10
 56 <210> SEQ ID NO: 2
 57 <211> LENGTH: 10
 58 <212> TYPE: PRT
 59 <213> ORGANISM: Homo sapiens
 61 <220> FEATURE:
 62 <221> NAME/KEY: PEPTIDE
 63 <222> LOCATION: (1)..(1)
 64 <223> OTHER INFORMATION: X=pyroglutamic acid
 67 <220> FEATURE:
 68 <221> NAME/KEY: PEPTIDE
 69 <222> LOCATION: (10)..(10)
 70 <223> OTHER INFORMATION: X=Gly-NH2
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```
                      RAW SEQUENCE LISTING              DATE: 06/05/2001
                      PATENT APPLICATION:  US/09/659,983B    TIME: 14:22:06

                      Input Set : A:\PTO.txt
                      Output Set: C:\CRF3\06052001\I659983B.raw


      73 <400> SEQUENCE: 2
      75 Xaa His Trp Ser His Gly Trp Tyr Pro Xaa
      76 1               5                   10
      78 <210> SEQ ID NO: 3
      79 <211> LENGTH: 20
      80 <212> TYPE: PRT
      81 <213> ORGANISM: artificial
      83 <220> FEATURE:
      84 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against
forms
      85        GnRH/ LHRH
      87 <220> FEATURE:
      88 <221> NAME/KEY: PEPTIDE
      89 <222> LOCATION: (1)..(1)
      90 <223> OTHER INFORMATION: X=pyroglutamic acid or Gln with attached tail of one or more
addi
      91        tional amino acid
      94 <220> FEATURE:
      95 <221> NAME/KEY: PEPTIDE
      96 <222> LOCATION: (3)..(3)
      97 <223> OTHER INFORMATION: X=Trp or N(indole)formyl-tryptophan
     100 <220> FEATURE:
     101 <221> NAME/KEY: PEPTIDE
     102 <222> LOCATION: (11)..(11)
     103 <223> OTHER INFORMATION: X=direct bond or a spacer group between Gly at position 10
and Gl
     104        n at position 1
     107 <220> FEATURE:
     108 <221> NAME/KEY: PEPTIDE
     109 <222> LOCATION: (13)..(13)
     110 <223> OTHER INFORMATION: X=Trp or N(indole)formyl-tryptophan
     113 <220> FEATURE:
     114 <221> NAME/KEY: PEPTIDE
     115 <222> LOCATION: (20)..(20)
     116 <223> OTHER INFORMATION: X=Gly-NH2 or Gly with attached tail of one or more amino
acids
     119 <220> FEATURE:
     120 <221> NAME/KEY: VARIANT
     121 <222> LOCATION: (10)..(20)
     122 <223> OTHER INFORMATION: variable repeat sequence <>10-20
     125 <400> SEQUENCE: 3
     127 Xaa His Xaa Ser Tyr Gly Leu Arg Pro Gly Xaa His Xaa Ser Tyr Gly
     128 1               5                   10                  15
     130 Leu Arg Pro Xaa
     131                 20
     133 <210> SEQ ID NO: 4
     134 <211> LENGTH: 21
     135 <212> TYPE: PRT
     136 <213> ORGANISM: artificial
     138 <220> FEATURE:
     139 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against
forms
     140        GnRH/ LHRH
```

[handwritten annotation: Xaa can only represent a single amino acid, nothing else]

142 <220> FEATURE:

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    143 <221> NAME/KEY: PEPTIDE
    144 <222> LOCATION: (1)..(1)
    145 <223> OTHER INFORMATION: X=pyroglutamic acid
    148 <220> FEATURE:
    149 <221> NAME/KEY: PEPTIDE
    150 <222> LOCATION: (6)..(6)
    151 <223> OTHER INFORMATION: X=D-Lys
    154 <220> FEATURE:
    155 <221> NAME/KEY: PEPTIDE
    156 <222> LOCATION: (11)..(11)
    157 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
    160 <220> FEATURE:
    161 <221> NAME/KEY: PEPTIDE
    162 <222> LOCATION: (16)..(16)
    163 <223> OTHER INFORMATION: X=D-Lys
    166 <220> FEATURE:
    167 <221> NAME/KEY: PEPTIDE
    168 <222> LOCATION: (21)..(21)
    169 <223> OTHER INFORMATION: X=Cys-NH2
    172 <400> SEQUENCE: 4
W--> 174 Xaa His Thr Ser Tyr Xaa Leu Arg Pro Gly Xaa His Thr Ser Tyr Xaa
    175 1               5                   10                  15
W--> 177 Leu Arg Pro Gly Xaa
    178              20
    180 <210> SEQ ID NO: 5
    181 <211> LENGTH: 21
    182 <212> TYPE: PRT
    183 <213> ORGANISM: artificial
    185 <220> FEATURE:
    186 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against
forms
    187       GnRH/ LHRH
    189 <220> FEATURE:
    190 <221> NAME/KEY: PEPTIDE
    191 <222> LOCATION: (1)..(1)
    192 <223> OTHER INFORMATION: X=pyroglutamic acid
    195 <220> FEATURE:
    196 <221> NAME/KEY: PEPTIDE
    197 <222> LOCATION: (4)..(4)
    198 <223> OTHER INFORMATION: X=amino acid substitution
    201 <220> FEATURE:
    202 <221> NAME/KEY: PEPTIDE
    203 <222> LOCATION: (6)..(6)
    204 <223> OTHER INFORMATION: X=D-Lys
    207 <220> FEATURE:
    208 <221> NAME/KEY: PEPTIDE
    209 <222> LOCATION: (11)..(11)
    210 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
    213 <220> FEATURE:
    214 <221> NAME/KEY: PEPTIDE
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215 <222> LOCATION: (14)..(14)
216 <223> OTHER INFORMATION: X=amino acid substitution
219 <220> FEATURE:
220 <221> NAME/KEY: PEPTIDE
221 <222> LOCATION: (16)..(16)
222 <223> OTHER INFORMATION: X=D-Lys
225 <220> FEATURE:
226 <221> NAME/KEY: PEPTIDE
227 <222> LOCATION: (21)..(21)
228 <223> OTHER INFORMATION: X=Cys-NH2
231 <400> SEQUENCE: 5
233 Xaa His Thr Xaa Tyr Xaa Leu Ala Pro Gly Xaa His Thr Xaa Tyr Xaa
234 1               5                   10                  15
236 Leu Arg Pro Gly Xaa
237             20
239 <210> SEQ ID NO: 6
240 <211> LENGTH: 21
241 <212> TYPE: PRT
242 <213> ORGANISM: artificial
244 <220> FEATURE:
245 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against
forms
246         GnRH/ LHRH
248 <220> FEATURE:
249 <221> NAME/KEY: PEPTIDE
250 <222> LOCATION: (1)..(1)
251 <223> OTHER INFORMATION: X=pyroglutamic acid
254 <220> FEATURE:
255 <221> NAME/KEY: PEPTIDE
256 <222> LOCATION: (6)..(6)
257 <223> OTHER INFORMATION: X=D-Lys
260 <220> FEATURE:
261 <221> NAME/KEY: PEPTIDE
262 <222> LOCATION: (8)..(8)
263 <223> OTHER INFORMATION: X=amino acid substitution
266 <220> FEATURE:
267 <221> NAME/KEY: PEPTIDE
268 <222> LOCATION: (11)..(11)
269 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
272 <220> FEATURE:
273 <221> NAME/KEY: PEPTIDE
274 <222> LOCATION: (16)..(16)
275 <223> OTHER INFORMATION: X=D-Lys
278 <220> FEATURE:
279 <221> NAME/KEY: PEPTIDE
280 <222> LOCATION: (18)..(18)
281 <223> OTHER INFORMATION: X=amino acid substitution
284 <220> FEATURE:
285 <221> NAME/KEY: PEPTIDE
286 <222> LOCATION: (21)..(21)
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Input Set : **A:\PTO.txt**
Output Set: **C:\CRF3\06052001\I659983B.raw**

```
287 <223> OTHER INFORMATION: X=Cys-NH2
290 <400> SEQUENCE: 6
292 Xaa His Thr Ser Tyr Xaa Leu Xaa Pro Gly Xaa His Thr Ser Tyr Xaa
293 1               5                    10             15
295 Leu Xaa Pro Gly Xaa
296              20
298 <210> SEQ ID NO: 7
299 <211> LENGTH: 21
300 <212> TYPE: PRT
301 <213> ORGANISM: artificial
303 <220> FEATURE:
304 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against
forms
305       GnRH/ LHRH
307 <220> FEATURE:
308 <221> NAME/KEY: PEPTIDE
309 <222> LOCATION: (1)..(1)
310 <223> OTHER INFORMATION: X=pyroglutamic acid
313 <220> FEATURE:
314 <221> NAME/KEY: PEPTIDE
315 <222> LOCATION: (6)..(6)
316 <223> OTHER INFORMATION: X=D-Lys
319 <220> FEATURE:
320 <221> NAME/KEY: PEPTIDE
321 <222> LOCATION: (10)..(10)
322 <223> OTHER INFORMATION: X=amino acid substutition
325 <220> FEATURE:
326 <221> NAME/KEY: PEPTIDE
327 <222> LOCATION: (11)..(11)
328 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
331 <220> FEATURE:
332 <221> NAME/KEY: PEPTIDE
333 <222> LOCATION: (16)..(16)
334 <223> OTHER INFORMATION: X=amino acid substitution
337 <220> FEATURE:
338 <221> NAME/KEY: PEPTIDE
339 <222> LOCATION: (20)..(20)
340 <223> OTHER INFORMATION: X=amino acid substitution
343 <220> FEATURE:
344 <221> NAME/KEY: PEPTIDE
345 <222> LOCATION: (21)..(21)
346 <223> OTHER INFORMATION: X=Cys-NH2
349 <400> SEQUENCE: 7
351 Xaa His Thr Ser Tyr Xaa Leu Arg Pro Xaa Xaa His Thr Ser Tyr Xaa
352 1               5                    10             15
354 Leu Arg Pro Xaa Xaa
355              20
357 <210> SEQ ID NO: 8
358 <211> LENGTH: 42
359 <212> TYPE: PRT
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to
<223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/659,983B**

DATE: 06/05/2001
TIME: 14:22:07

Input Set : **A:\PTO.txt**
Output Set: **C:\CRF3\06052001\I659983B.raw**

```
L:13  M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:53  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:75  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:81  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:136 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:183 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:242 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:301 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:360 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:434 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:481 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:519 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:528 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:587 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:640 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:646 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:693 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
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